S1. SNP retention of rAAV gene targeting colonies in parental HCT116 cells.

									_egends:	Viral	Genomic
Ndel	EcoRI	LHP	Ncol	Asel	Sspl	Sacl	RHP	Xbal	Sbfl	#	%
+	+	+	+	+	+	+	-	-	-	5	2.2%
+	+	+	+	+	+	-	-	-	-	1	0.4%
+	+	+	+	+	-	+	-	-	+	1	0.4%
-	+	+	+	+	+	+	+	+	-	2	0.9%
-	+	+	+	+	+	+	+	-	-	14	6.1%
-	+	+	+	+	+	+	-	+	-	1	0.4%
-	+	+	+	+	+	+	-	-	-	9	3.9%
-	+	+	+	+	+	-	-	-	+	2	0.9%
-	+	+	+	+	+	-	-	-	-	17	7.4%
-	+	+	+	+	-	+	-	-	-	7	0.4%
-	+	+	+	+	-	-	-	-	-		3.0%
-	+	+	-	+	+	-	-	-	-	2	0.9% 0.9%
-	+	-	+	+	+	+	-	-	-		
-	+	-	+	+	-	+	+	+	-	1	0.4% 0.4%
	-								-	3	1.3%
-	-	+	+	+	+	+	+	+	-	3	1.3%
_	-	+	+	+	+	+	-	-	_	6	2.6%
	-	+	+	+	+	-	+	-	-	2	0.9%
_	-	+	+	+	+	-	-	-	_	13	5.7%
_	_	+	+	+	-	-	_	-	_	4	1.7%
_	-	+	-	+	+	+	+	+	-	11	4.8%
-	-	+	-	+	+	+	+	-	-	5	2.2%
-	-	+	-	+	+	-	+	+	_	2	0.9%
-	-	+	-	+	+	-	-	-	-	1	0.4%
-	-	+	-	+	-	+	+	-	-	1	0.4%
-	-	+	-	+	-	-	+	-	-	1	0.4%
-	-	+	-	-	-	+	+	+	-	1	0.4%
-	-	-	+	+	+	+	+	+	+	2	0.9%
-	-	•	+	+	+	+	+	+	-	3	1.3%
-	1	1	+	+	+	+	+	-	-	1	0.4%
-	-	-	+	+	+	+	-	-	-	3	1.3%
-	-	-	+	+	+	-	+	-	-	1	0.4%
-	-	-	+	+	+	-	-	-	-	1	0.4%
-	-	-	-	+	+	+	+	+	+	5	2.2%
-	-	-	-	+	+	+	+	+	-	35	15.2%
-	-	-	-	+	+	+	+	-	-	16	7.0%
-	-	-	-	+	+	+	-	+	+	3	1.3%
-	-	-	-	+	+	+	-	+	-	2	0.9%
-	-	-	-	+	+	+	-	-	-	6	2.6%
-	-	-	-	+	+	-	+	-	-	2	0.9%
-	-	-	-	+	+	-	-	-	-	19	8.3%
-	-	-	-	+	-	+	+	+	-	2	0.9%
-	-	-	-	+	-	-	-	-	-	2	0.9%
-	-	-	-	-	+	+	+	+	-	3	1.3%
-	-	-	-	-	+	+	-	-	-	4	0.4% 1.7%
7			105	220		148	116	76		230	
_	66	115			209				13	23 0	100.00%
3.04%	28.70%	50.00%	45.65%	95.65%	90.87%	64.35%	50.43%	33.04%	5.65%		